# caArray Use Cases

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### Use case 1: Import Microarray dataset as MAGE-ML

We would like to use caArray to import dataset in MAGE-ML format to an in-house MAGE-OM compliant Oracle database.

# Use case 2: Export Microarray dataset as MAGE-ML

We would like to use caArray to export MAGE-ML so that we can export any data stored in an in-house MAGE-OM compliant Oracle database to any other MAGE-OM compliant databases.

### **Use case 3: Upload Affymetrix Microarray**

We would like to use caArray to upload the MIAME information along with the following affymetrix microarray files to an in-house MAGE-OM compliant Oracle database assuming 2 treatments and 4 replicates per treatment for the experiment to be uploaded. In addition, we would like to have caArray to store .CEL files in a designated file system directory for R routines to use.

```
EXP1 TRT1 REP1.CHP
EXP1 TRT1 REP2.CHP
EXP1_TRT1_REP3.CHP
EXP1 TRT1 REP4.CHP
EXP1_TRT2_REP1.CHP
EXP1 TRT2 REP2.CHP
EXP1 TRT2 REP3.CHP
EXP1_TRT2_REP4.CHP
EXP1_TRT1_REP1.CEL
EXP1 TRT1 REP2.CEL
EXP1 TRT1 REP3.CEL
EXP1_TRT1_REP4.CEL
EXP1 TRT2 REP1.CEL
EXP1_TRT2_REP2.CEL
EXP1 TRT2 REP3.CEL
EXP1_TRT2_REP4.CEL
EXP1 TRT1 REP1.rpt
EXP1 TRT1 REP2.rpt
EXP1 TRT1 REP3.rpt
EXP1_TRT1_REP4.rpt
```

```
EXP1_TRT2_REP1.rpt
EXP1_TRT2_REP2.rpt
EXP1_TRT2_REP3.rpt
EXP1_TRT2_REP4.rpt

EXP1_TRT1_REP1.exp
EXP1_TRT1_REP2.exp
EXP1_TRT1_REP3.exp
EXP1_TRT1_REP4.exp
EXP1_TRT2_REP1.exp
EXP1_TRT2_REP1.exp
EXP1_TRT2_REP2.exp
EXP1_TRT2_REP2.exp
EXP1_TRT2_REP3.exp
EXP1_TRT2_REP3.exp
EXP1_TRT2_REP4.exp
```

In addition, we would like to use caArray to upload the following image files in a designated file system directory along with the experiment and store the URL in the Oracle database for future retrieval, viewing and image reanalyzing when new image analyzing software becomes available.

```
EXP1_TRT1_REP1.DAT
EXP1_TRT1_REP2.DAT
EXP1_TRT1_REP3.DAT
EXP1_TRT1_REP4.DAT
EXP1_TRT2_REP1.DAT
EXP1_TRT2_REP2.DAT
EXP1_TRT2_REP3.DAT
EXP1_TRT2_REP4.DAT
EXP1_TRT2_REP4.DAT
```

## **Use case 4: Upload 2-channel Microarray**

We would like to use caArray to upload the MIAME information along with the following 2-channel microarray files to the backend Oracle database assuming 2 treatments and 4 replicates per treatment for the experiment to be uploaded.

```
EXP1_TRT1_REP1.GPR
EXP1_TRT1_REP2.GPR
EXP1_TRT1_REP3.GPR
EXP1_TRT1_REP4.GPR
EXP1_TRT2_REP1.GPR
EXP1_TRT2_REP2.GPR
EXP1_TRT2_REP3.GPR
EXP1_TRT2_REP4.GPR
EXP1_TRT1_REP1.GAL
EXP1_TRT1_REP2.GAL
EXP1_TRT1_REP3.GAL
```

EXP1\_TRT1\_REP4.GAL EXP1\_TRT2\_REP1.GAL EXP1\_TRT2\_REP2.GAL EXP1\_TRT2\_REP3.GAL EXP1\_TRT2\_REP4.GAL

With both use cases 3 and 4, we would like to use caArray to assign the initial ownership of the uploaded experiment so that only the owner can view the experiment until he/she authorize others to view/analyze the datasets in a separate module. This owner may or may not be the user uploading the experiment.

### **Relevant issues for Architecture:**

- 1. MAGE-OM compliant
- 2. Ability to upload large datasets in a reasonable speed
- 3. Ability to generate MAGE-ML and import MAGE-ML in a reasonable speed
- 4. Ability to track MIAME such as experiment condition, experiment design, biosource, array design, sample extraction condition, hybridization condition and scanning parameters et al
- 5. Ability to track ownership